

Amendments to the Specification:

Please replace the text at page 24, line 10 with the following amended text:

regulatory system from Valentis (~~www.geneswitch.com~~), sold by Invitrogen (catalog

Please replace the text at page 24, line 29 with the following amended text:

PPLLQAIELFDHQGAEPALFLGLELIICGLEKQLKCESGS (SEQ ID NO: 18); (a-1) **tTA**. (TetR-VP16)

Please replace the text at page 24, line 37 with the following amended text:

YGG (SEQ ID NO: 19); (a-2) **tTA2** (TetR-FFF) (Baron et al., 1997):

Please replace the text at page 25, line 9 with the following amended text:

ALDDFDLMDLPADALDDFDLMDLPG (SEQ. ID. No.: 20); (a-3) **tTA-p65** (TetR-p65) (Urlinger et al.,

Please replace the text at page 25, line 19 with the following amended text:

GLPNGLSGDEDFSSIADMDFSALLSQISS (SEQ. ID. No.: 21); (b) **Lac repressor**: GenBank

Please replace the text at page 25, line 27 with the following amended text:

TASPRALADSLMQLARQVSRLESGQ (SEQ. ID. NO.: 22); (c) **Xylose repressor**: GenBank accession

Please replace the text at page 25, line 35 with the following amended text:

LDMITM (SEQ. ID. NO.: 23);

Please replace the text at page 26, line 11 with the following amended text:

GCEEKVNDVAVKLS (SEQ. ID. NO.: 24); (e) TetR-based T-Rex system: the TetR sequence is

Please replace the text at page 26, line 17 with the following amended text:

VLAQIAAILCLMFPEHDDFQLLQAHA (SEQ. ID. NO.: 25); (g) ***Streptomyces coelicolor***

Please replace the text at page 26, line 23 with the following amended text:

EDAQDIMAARGGDTVAEMLDLDRDFEFALDLLVAGIDAMVEQA (SEQ. ID. NO.: 26); (h) ***Streptomyces***

Please replace the text at page 26, line 29 with the following amended text:

DYQDLEHRYALLQKHILPAIAVPSVLAALDLSEERGARLAAELAPTGKD (SEQ. ID. NO.: 27); (i) **TraR-**

Please replace the text at page 27, line 5 through page 28, line 31, with the following amended text:

QHRHITAVTNYHRQWQSTYFDKKFEALDPVVKRARSRKHIFTWSGEHERPTLSKD
 ERAFYDHASDFGIRSGITIPKTANGFMSMFTMASDKPVIDLDREIDAVAAAATIGQI
 HARISFLRTTPTAEDAACVDPKEATYLRWIAVGKTMEEIADVEGVKYNVSVRVKLRE
 RMKRFDVRSKAHLTALAIRRKLI (SEQ. ID. NO. 28); (j-i) The skilled person knows that almost any
 fusion of the Phy sequences to a Gal4-DBD (aa's 1-63, 1-95, 1-141) would be
 active. (j-ii) **Gal4-hpr-p65** from pSwitch (Invitrogen: Geneswitch_man.pdf):
 MDSQQPDLKLLSSIEQACDICRLKKLKCSKEPKCAKCLKNNWECRYSPKTKRSPL
 TRAHLTEVESRLERLEQLFLLIFPREDLDMILKMDSLQDIKALLEFPGVDQKKFNKV
 RVVRALDAVALPQPVGVNPNESQALSQRFTFSPGQDIQLIPPLINLLMSIEPDVIYAG
 HDNTKPDTSLSLTSLNQLGERQLLSVVKWSKSLPGFRNLHIDDQITLIQYSWMSL
 MVFGLGWRSYKHVSGQMLYFAPDLILNEQRMKESSFYSLCLTMWQ1PQEFVKLQV
 SQEEFLCMKVLLLLNTIPLEGLRSQTQFEEMRSSYIRELIKAIGLRQKGVVSSSQRF
 YQLTKLLDNLDLVKQLHLYCLNTFIQSRALSVEFPEMMSEVIAGSTPMEFYLPDT
 DDRHRIEEKRKRTYETFKSIMKKSPFSGPTDPRPPPRRIAVPSRSSASVPKPAPQP
 YPFTSSLSTINYDEFPTMVFPSPGQISQASÄLAPAPPQVLPQAPAPAPAPAMVSALA
 QAPAPVPVLAPGPPQAVAPPAPKPTQAGEGTLSEALLQLQFDDDLGALLGNSTD
 PAVFTDLASVDNSEFQQLLNQGIPVAPHTTEPMLMEYPEAITRLVTGAQRPPDPAP
 APLGAPGLPNGLSGDEDFSSSIADMDFSALLSQISS (SEQ. ID. NO.: 29); (j-iii-1) ZHFD1-FKBP fusion
 (www.ariad.com/regulationkits)
 MDYPAAKRVKLDSESRERPYACPVESCDRRFSRDELTRHIRIHTGQKPFQCRICMR
 NFSRSDHLTTHIRTHTGGRRRKRKRTSIETNIRVALEKSFLNQKPTSEEITMIADQL
 NMEKEVIRVWFCNRRQKEKRINTRGVQVETISPGDGRTFPRGQTCWHYTGMLE
 DGKKFDSSDRNKPFFKMLGKQEVIRGWEEGVAQMSVGQRAKLTISPDYAYGAT
 GHPGIIPPHATLVFDVELLKLEVEGVQVETISPGDGRTFPRGQTCWHYTGMLED
 GKKFDSSDRNKPFFKMLGKQEVIRGWEEGVAQMSVGQRAKLTISPDYAYGATG
 HPGIIPPHATLVFDVELLKLETRGVQVETISPGDGRTFPRGQTCWHYTGMLEDG
 KKFDSSDRNKPFFKMLGKQEVIRGWEEGVAQMSVGQRAKLTISPDYAYGATGH
 PGIIPPHATLVFDVELLKLETSY (SEQ. ID. NO.: 30); (j-iii-2) FRB-p65 fusion
 (www.ariad.com/regulationkits):
 MDYPAAKRVKLDSESRILWHEMWHEGLEEASRLYFGERNVKGMFEVLEPLHAMMER
 GPQTLKETSFNQAYGRDLMEAEQWCRKYMKSGNVKDLLQAWDLYYHVFRISK
 RDEFPTMVFPSPGQISQASALAPAPPQVLPQAPAPAPAPAMVSALAQAPAPVPVLA
 PGPQAVAPPAPKPTQAGEGTLSEALLQLQFDDDLGALLGNSTDPAVFTDLASV
 DNSEFQQLLNQGIPVAPHTTEPMLMEYPEAITRLVTGAQRPPDPAPAPLGAPGLPN
 GLLSGDEDFSSSIADMDFSALLSQISSTSY (SEQ. ID. NO.: 31); (j-iv-1) VgEcR from pVgRXR
 (<http://www.invitrogen.com/content/sfs/vectors/pvgxr.pdf>):

MAPPTDVSLGDELHLDGEDVAMAHADALDDFDLMLGDGDSPPGFTPHDSAPY
GALDMADFEFEQMFTDALGIDEYGGKLLGTSRRISNSISSGRDDLSPSSSLNGYSA
NESCDAKKSKKGPAPRVQEELCLVCGDRASGYHYNALTCGSCKVFFRRSVTKSA
VYCKKFRACEMDMYMRRKCQECRLKKCLAVGMRPECWPENQCAMKRREEKA
QKEKDKMTTSPSSQHGGNGSLASGGGQDFVKKEILDLMTCPEPPQHATIPLLPDEIL
AKCQARNIPSLTYNQLAVIYKLIWYQDGYEQPSEEDLRRIMSQPDENESQTDVSFR
HITEITILTVQLIVEFAKGLPAFTKIPQEDQITLLKACSSEVMMLRMARRYDHSSDSIF
FANNRSYTRDSYKMAGMADNIEDLLHFCRQMFMSMKVDNVEYALLTAIVIFSDRPGL
EKAQLVEAIQSYIDTLRIYILNRHCGDSMSLVFYAKLLSILTELRTLGNAEMCFS
LKLKNRKLPKFLEEIWDVHAIPPSVQSHLQITQEENERLERAERMRAVGGAITAGI
DCDSASTSAAAAAAQHQPQPQPQPSSLTQNDQHQQTQPQLQPQLPPQLQGQ
LQPQLQPQLQTQLQPQIQPQPQLLPVSAPVPASVTAPGSLSAVSTSSEYMGGSA
IGPITPATTSSITA AVTASSTTSAPVPMGNGVGVGVGVGGNVSMYANAQTAMALMG
VALHSHQEQLIGGVAVKSEHSTTA (SEQ. ID. NO.: 32); (j-iv-2) RXR from pVgRXR
(<http://www.invitrogen.com/content/cfs/vectors/pvgrr.pdf>);
MDTKHFLPLDFSTQVNSSLTPTGRGSMAAPSLHPSLGPGLGSPGQLHSPISTLSS
PINGMGPPFSVISSPMGPHSMSVPTTPTLGFSTGSPQLSSPMNPVSSSEDIKPLG
LNGVLKVAHPSGNMASTFKHICAICGDRSSGKHGYVYSCEGCKGFFKRTVRKDL
TYTCRDNKDCLIDKRQRNRCQYCRYQMCLAMGMKREAVQEERQRGKDRNENEV
ESTSSANEDVPVERILEAEALAVEPKTETYVEANVGLNPSSPNDPVTNICQAADKQL
FTLVEWAKRIPHFSELPLDDQVILLRAGWNELLASFHSRSIAVKDGI LLATGLHVHR
NSAHSAGVGAIFDRVLTELVS KM RDMQMDKTELGLRAIVLFNPDSKGLSNPAEV
EALREKVYASLEAYCKHKYPEQPGRFAKLLRLPALRSIGLKCLEHLFFFKLIGDTPI
DTFLMEMLEAPHQMT (SEQ. ID. NO.: 33).